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<213> homo sapiens

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77-7	т1 -	C1:-	Λ ~	т1 -	Ι	7\ ~~ ~~	ui-	D~~	C1	τ <i>τ</i> ~ 1	Luc	₹7 ¬ ¬	C1~	C^~	Lva
			Asn 420			_		425			_		430		
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Lys 465	Val	Lys	Arg	Ser	Leu 470	Glu	Asp	Glu	Pro	Arg 475	Leu	Val	Leu	Trp	Al,a 480
Leu	Leu	Val	Тут	Asn 485	Ala	Thr	Asn	Asn	Thr 490	Asn	Leu	Glu	Gly	Lys 495	Ile
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Arg	Leu	His 515	Thr	Val	Asn	Val	Arg 520	Gln	Leu	Gly	His	Cys 525	Leu	Ala	Met
Glu	Glu 530		L;/s	Gly	Tyrr	Tyr 535		Pro	Ser	Ile	Gln 540	Pro	Ser	Glu	Тут
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Thr E	Phe	Ile	Ser	Туг		Gly	Суѕ	Gly	Ile		Ala	Ile	Phe	Ser	
865	-1			m)	870	** 7		D1 .	0.1	875	T	3	7	7	880
Ala T				885					890					895	
Pro S	Ser	Lys	Ile 900	Leu	Met	Asn	Leu	Ser 905	Thr	Ala	Leu	Leu	Phe 910	Leu	Asn
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Lys V	/al	Phe	Asn	Thr		Tle	Ara	Ara	Tvr		Leu	Lvs	Phe	Cvs	
י פינים	v a i	1110	11011	965	1 1 1	110	9	9	970	110	200	272		975	
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Leu C			T ON	Dho	Tlo			Dho	ціс	C: 15			Lvc	Glu	Agn
11.05	3 T I I	GIY	гел	rne	1110		TIE	FILE	1115	111!		Mec	CYL	Giu	1120
Val C	aln	Lvs	Gln	Trp			His	Leu	Cvs			Ara	Phe	Ara	
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<212> PRT

<213> homo sapiens

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Ile	Pro	Glu	Leu 180	Ser	Ala	Phe	Thr	Leu 185	СЛа	Phe	Glu	Ala	Thr 190	Lys	Val
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	Thr	275					280					235			
-	390 FA2				_	295					300				
305	Asp				310					315					320
	Leu			325		_			330					335	
	Asn		340					345					350		
	Leu	355					360					365			
	Ser 370					375					380				
385	Ser				390					395					400
	Ile			405					410					415	
	Ile		420					425					430		
	Ala	435	_				440					445			
	Val 450					455					460				
465	Val	-			470		_			475				_	480
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	Gln		500					505					510		
	Leu	515					520					525			
	Glu 530					535					540				
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rne	тАг	Abli	MIG	565	nsli	LIO	ьeu	va⊥	570	тАт	ттЪ	GTÅ	FIO	575	qan

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Ile	Asp	Glu	Leu 660	Ala	Phe	Lys	Ile	Asp 665	Leu	Asn	Ser	Thr	Ser 670	His	Val
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<213> homo sapiens

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<213> homo sapiens

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Gly	His	Glu 195	Asp	Ser	Asp	Trp	Thr 200	Ala	Phe	Ser	Tyr	Ser 205	Asn	Ala	Ser
Phe	Thr 210	Gln	Leu	Leu	Ser	Phe 215	Gly	Lys	Ala	Lys	Ser 220	Gly	Tyr	Phe	Leu
Ser 225	Ile	Ser	Asp	Ser	Lys 230	Cys	Leu	Leu	Asn	Asn 235	Ala	Leu	Pro	Val	Lys 2∴0
Glu	Lys	Glu	Asp	Ile 245	Phe	Ala	Glu	Ser	Phe 250	Glu	Gln	Leu	Cys	Leu 255	Vāl
Trp	Asn	Asn	Ser 250		Gly	Ser	Ile	Gly 265		Asn	Phe	Lys	Arg 270	Asn	Туг
Glu	Thr	Val 275	Pro	Cys	Asp	Ser	Thr 230	Ile	Ser	L7.2	Vāl	Ile 285	Pr.o	Gly	Asn
Gly	Lys 290	Leu	Leu	Leu	Gly	Ser 295	Asn	Gln	Asn	Glu	Ile 300	Val	Ser	Leu	Lys
Gly 305	Asp	Ile	Tyr	Asn	Phe 310	Arg	Leu	Trp	Asn	Phe 315	Thr	Met	Asn	Ala	Lys 320
He	Leu	Ser	Asn	Leu 325	Ser	Cys	Asn	Val	Lys 330	Gly	Asn	Val	Vāl	Asp 335	Trp
Gln	Asn	Asp	Phe 340	Trp	Asn	Ile	Pro	Asn 345	Leu	Ala	Leu	Lys	Ala 350	Glu	Ser
Asn	Leu	Ser 355	Cys	Gly	Ser	Tyr	Leu 360	Ile	Pro	Leu	Pro	Ala 365	Ala	Glu	Leu
Ala	Ser 370	Cys	Ala	Asp	Leu	Gly 375	Thr	Leu	Cys	Gln	Ala 380	Thr	Val	Asn	Ser
Pro 385	Ser	Thr	Thr	Pro	Pro 390	Thr	Val	Thr	Thr	Asn 395	Met	Pro	Vāl	'Thr	Asn 400
Arg	Ile	Asp	Lys	Gln 405	Arg	Asn	Asp	Gly	Ile 410	Ile	Tyr	Arg	Ile	Ser 415	Val
Val	Ile	Gln	Asn 4.10	Ile	Leu	Arg	His	Pro 425	Glu	Val	Lys	Val	Gln 430	Ser	Lys
Val	Ala	Glu 435	Trp	Leu	Asn	Ser	Thr 440	Phe	Gln	Asn	Trp	Asn 445	Tyr	'Thr	Vāl
Туг	Val 450	Val	Asn	Ile	Ser	Phe 455	His	Leu	Ser	Ala	Gly 460	Glu	Asp	Lys	Ile
Lуs 465	Val	Lys	A:-g	Ser	Leu 470	Glu	Asp	Glu	Pro	Arg 475	Leu	Val	Leu	Trp	Ala 480
Leu	Leu	Val	Tyr	Asn 485	Ala	Thr	Asn	Asn	Thr 490	Asn	Leu	Glu	Gly	Lys 495	Ile
Ile	Gln				Leu										Leu
Arg	Leu	His 515	Thr	Val	Asn	Val	Arg 520	Gln	Leu	Gly	His	Cys 525	Leu	Ala	Met
Glu	Glu 530	Pro	L;/s	Gly	Тут	Tyr 535	Trp	Pro	Ser	Ile	Gln 540	Pro	Ser	Glu	Tyr
Val 545	Leu	Pro	Cys	Pro	Asp 550	Lys	Pro	Gly	Phe	Ser 555	Ala	Ser	Ar-g	Ile	Cys 560
Phe	Tyr	Asn	Ala	Thr 565	Asn	Pro	Leu	Val	Thr 570	Туг	Trp	Gly	Pro	Val 575	Asp
Ile	Ser	Asn	Cys 560	Leu	Lys	Glu	Ala	Asn 585	Glu	Val	Ala	Asn	Gln 590	Ile	Leu
Asn	Leu	Thr 595	Ala	Asp	Gly	Gln	Asn 600	Leu	Thr	Ser	Ala	Asn 605	Ile	Thr	Asn
Ile	Val 610	Glu	Gln	Val	Lys	Arg 615	Ile	Val	Asn	Lys	Glu 620	Glu	Asn	Ile	Asp
Ile 625	Thr	Leu	Gly	Ser	Thr 630	Leu	Met	Asn	Ile	Phe 635	Ser	Asn	Ile	Leu	Ser 640

Ser	Ser	Asp	Ser	Asp 645	Leu	Leu	Glu	Ser	Ser 650	Ser	Glu	Ala	Leu	Lys 655	Thr
Ile	Asp	Glu	Leu 650	Ala	Phe	Lys	Ile	Asp 665	Leu	Asn	Ser	Thr	Ser 670	His	Val
Asn	Ile	Thr 675	Thr	Arg	Asn	Leu	Ala 680	Leu	Ser	Val	Ser	Ser 635	Leu	Leu	Pro
Gly	Thr 690	Asn	Ala	Ile	Ser	Asn 695	Phe	Ser	Ile	Gly	Leu 700	Pro	Ser	Asn	Asn
Glu 705	Ser	Tyr	Phe	Gln	Met 710	Asp	Phe	Glu	Ser	Gly 715	Gln	Val	Asp	Pro	Leu 720
Ala	Ser	Val	Ile	Leu 725	Pro	Pro	Asn	Leu	Leu 730	Glu	Asn	Leu	Ser	Pro 735	Glu
Asp	Ser	Val	Leu 740	Val	Arg	Arg	Ala	Gln 745	Phe	Thr	Phe	Phe	Asn 750	Lys	Thr
Gly	Leu	Phe 755	Gln	Asp	Val	Gly	Pro 750	Gln	Arg	Lys	Thr	Leu 765	Val	Ser	Tyr
	770		Cys			775					780				
Pro 785	Val	Gln	Ile	Lys	Ile 790	Lys	His	Thr	Arg	Thr 795	Gln	Glu	Val	His	His 800
Pro	Ile	C7.2	Ala	Phe 305	Trp	Asp	Leu	Asn	Lys 810	Asn	Lys	Ser	Phe	Gly 815	Gly
Trp	Asn	Thr	Ser 820	Gly	Cys	Va.	Ala	His 825	Arg	Asp	Ser	Asp	Ala 830	Ser	Glu
Thr	Val	835 67.s	Leu	Cys	Asn	His	Phe 840	Thr	His	Phe	Gly	Val 845	Leu	Met	Asp
	850		Ser			855					860				
Thr 365	Phe	Ile	Ser	Туг	Ile 870	Gly	Cys	Gly	Ile	Ser 875	Ala	Ile	Phe	Ser	Ala 880
Ala	Thr	Leu	Leu	Thr 385	Tyr	Val	Ala	Phe	Glu 890	Lys	Leu	Arg	Arg	Asp 895	Tyr
			Ile 900					905					910		
Leu	Leu	Phe 915	Leu	Leu	Asp	Gly	Trp 920	Ile	Thr	Ser	Phe	Asn 925	Val	Asp	Gly
Leu	Cys 930	Ile	Ala	Val	Ala	Val 935	Leu	Leu	His	Phe	Phe 940	Leu	Leu	Ala	Thr
945		_		_	950					955	_				960
Lys	Val	Phe	Asn	Thr 965	Tyr	Ile	Arg	Arg	Tyr 970	Ile	Leu	Lys	Phe	Cys 975	Ile
			Gly 980					985					990		
_		995	Asn			_	1000)				1005	5		
	1010)	Phe			1015	5				1020)			
		Gly	Tyr	Phe			Met	Phe	Phe			Ile	Ala	Met	
1025 Ile		Val	Met	Val	1030 Gln		Cws	Glv	Ara	1035 Asn		Lvs	Ara	Ser	1040 Asn
			Arg	1045	5				1050)				1055	5
_			1060)				1065	5				1070)	
ьeu	1111	1075	Leu	ьeu	GTÀ	мес	1080		,aΤ7,	rne	AId	1085		Ala	ттħ

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75

80

70

Ile	Lys	His	Thr	Arg 85	Thr	Gln	Glu	Val	His 90	His	Pro	Ile	Cys	Ala 95	Phe
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Cys	Val	Ala 115	His	Arg	Asp	Ser	Asp 120	Ala	Ser	Glu	Thr	Val 125	Cys	Leu	Cys
Asn	His 130	Phe	Thr	His	Phe	Gly 135	Val	Leu	Met	Asp	Leu 140	Pro	Arg	Ser	Ala
Ser 145	Gln	Leu	Asp	Ala	Arg 150	Asn	Thr	Lys	Val	Leu 155	Thr	Phe	Ile	Ser	Tyr 160
lle	Gly	Cys	Gly	Ile 165	Ser	Ala	Ile	Phe	Ser 170	Alā	Ala	Thr	Leu	Leu 175	Thr
Тут	Val	Ala	Phe 130	Glu	Lys	Leu	Arg	Arg 135	Asp	Tyr	Pro	Ser	Lys 190	Ile	Leu
Met	Asn	Leu 195	Ser	Thr	Ala	Leu	Leu 200	Phe	Leu	Asn	Leu	Leu 205	Phe	Leu	Leu
Asp	Gly 210	Trp	Ile	Thr	Ser	Phe 215	Asn	Val	Asp	Glу	Leu 220	Суѕ	Ile	Ala	Val.
225					230					235				Met	240
Leu	Glu	Ala	Ile	Hıs 245	Met	Tyr	Ile	Ala	Leu 250	Val	Lys	Val	Phe	Asn 255	Thr
Tyr	Ile	Arg	Arg 250	Tyr	Ile	Leu	ГЛЯ	Phe 265	Cys	Ile	Ile	Gly	Trp 270	Gly	Leu
Pro	Ala	Leu 275	Val	Val	Ser	Val	Val 280	Leu	Ala	Ser	Arg	Asn 285	Asn	Asn	Glu
Val	_	Gly	Lys	Glu	Ser		Gly	Lys	Glu	Lys		Asp	Glu	Phe	Cys
	290					295					300				
3 (* 5	Ile				310	Ile				315	Cys			Tyr	320
305 Gly	Ile Val	Met	Phe	Phe 325	310 Leu	Ile Asn	Ile	Ala	Met 330	315 Phe	Cys Ile	Val	Val	Met 335	320 Val
305 Gly Gln	Ile Val Ile	Met Cys	Phe Gly 340	Phe 325 Arg	310 Leu Asn	Ile Asn Gly	Ile Lys	Ala Arg 345	Met 330 Ser	315 Phe Asn	Cys Ile Arg	Val Thr	Val Leu 350	Met 335 Arg	320 Val Glu
305 Gly Gln Glu	Ile Val Ile Val	Met Cys Leu 355	Phe Gly 340 Arg	Phe 325 Arg Asn	310 Leu Asn Leu	Ile Asn Gly Arg	Ile Lys Ser 360	Ala Arg 345 Val	Met 330 Ser Val	315 Phe Asn Ser	Cys Ile Arg Leu	Val Thr Thr 365	Val Leu 350 Phe	Met 335 Arg Leu	320 Val Glu Leu
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Gly Glu Gly Gly Pro 385	Ile Val Ile Val Met 370 Phe	Met Cys Leu 355 Thr	Phe Gly 340 Arg Trp	Phe 325 Arg Asn Gly Leu	310 Leu Asn Leu Phe 390	Asn Gly Arg Ala 375 Ser	Ile Lys Ser 360 Phe	Ala Arg 345 Val Phe	Met 330 Ser Val Ala Asn	315 Phe Asn Ser Trp Ser 395	Cys Ile Arg Leu Gly 380 Leu	Val Thr Thr 365 Pro	Val Leu 350 Phe Leu Gly	Met 335 Arg Leu Asn	320 Val Glu Leu Ile Phe 400
Gly Glu Gly Pro 385 Ile	Ile Val Ile Val Met 370 Phe	Met Cys Leu 355 Thr Met	Phe Gly 340 Arg Trp Tyr	Phe 325 Arg Asn Gly Leu His 405	310 Leu Asn Leu Phe 390 Cys	Asn Gly Arg Ala 375 Ser Ala	Ile Lys Ser 360 Phe Ile	Ala Arg 345 Val Phe Phe	Met 330 Ser Val Ala Asn Glu 410	315 Phe Asn Ser Trp Ser 395 Asn	Cys Ile Arg Leu Gly 380 Leu Val	Val Thr Thr 365 Pro Gln Gln	Val Leu 350 Phe Leu Gly	Met 335 Arg Leu Asn Leu Gln 415	320 Val Glu Leu Ile Phe 400 Trp
Gly Glu Gly Pro 385 Ile Arg	Ile Val Ile Val Met 370 Phe Phe Arg	Met Cys Leu 355 Thr Met Ile	Phe Gly 340 Arg Trp Tyr Phe Leu 420	Phe 325 Arg Asn Gly Leu His 405 Cys	Asn Leu Phe Phe 390 Cys	Asn Gly Arg Ala 375 Ser Ala Gly	Ile Lys Ser 360 Phe Ile Met	Ala Arg 345 Val Phe Phe Lys Phe 425	Met 330 Ser Val Ala Asn Glu 410 Arg	315 Phe Asn Ser Trp Ser 395 Asn Leu	Cys Ile Arg Leu Gly 380 Leu Val Ala	Val Thr Thr 365 Pro Gln Gln	Val Leu 350 Phe Leu Gly Lys Asn 430	Met 335 Arg Leu Asn Leu Gln 415 Ser	320 Val Glu Leu Ile Phe 400 Trp
Gly Glu Gly Pro 385 Ile Arg	Ile Val Ile Val Met 370 Phe Phe Arg Ser	Met Cys Leu 355 Thr Met Ile His Lys 435	Phe Gly 340 Arg Trp Tyr Phe Leu 420 Thr	Phe 325 Arg Asn Gly Leu His 405 Cys	310 Leu Asn Leu Phe 390 Cys Cys	Asn Gly Arg Ala 375 Ser Ala Gly Asn	Ile Lys Ser 360 Phe Ile Met Arg Ile 440	Ala Arg 345 Val Phe Phe Lys Phe 405 Ile	Met 330 Ser Val Ala Asn Glu 410 Arg	315 Phe Asn Ser Trp Ser 395 Asn Leu Lys	Cys Ile Arg Leu Gly 380 Leu Val Ala Ser	Val Thr Thr 365 Pro Gln Gln Asp Ser 445	Val Leu 350 Phe Leu Gly Lys Asn 430 Asp	Met 335 Arg Leu Asn Leu Gln 415 Ser	320 Val Glu Leu Ile Phe 400 Trp Asp Leu
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Gly Gln Glu Gly Pro 385 Ile Arg Trp Gly Thr 465	Ile Val Ile Val Met 370 Phe Phe Arg Ser Lys 450 Ser	Met Cys Leu 355 Thr Met Ile His Lys 435 Ser Lys	Phe Gly 340 Arg Trp Tyr Phe Leu 420 Thr Leu Ser	Phe 325 Arg Asn Gly Leu His 405 Cys Ala Ser Lys	Asn Leu Phe 390 Cys Cys Thr Ser 470	Asn Gly Arg Ala 375 Ser Ala Gly Asn Ser 455 Ser	Ile Lys Ser 360 Phe Ile Met Arg Ile 440 Ser	Ala Arg 345 Val Phe Phe Lys Phe 425 Ile Ile	Met 330 Ser Val Ala Asn Glu 410 Arg Lys Gly Thr	Asn Ser Trp Ser 395 Asn Leu Lys Ser Tyr 475	Cys Ile Arg Leu Gly 380 Leu Val Ala Ser Asn 460 Phe	Val Thr Thr 365 Pro Gln Gln Asp Ser 445 Ser Lys	Val Leu 350 Phe Leu Gly Lys Asn 430 Asp Thr	Met 335 Arg Leu Asn Leu Gln 415 Ser Asn Tyr	320 Val Glu Leu Ile Phe 400 Trp Asp Leu Leu Ser 480
Gly Gln Glu Gly Pro 385 Ile Arg Trp Gly Thr 465 His	Ile Val Ile Val Met 370 Phe Phe Arg Ser Lys 450 Ser Thr	Met Cys Leu 355 Thr Met Ile His Lys 435 Ser Lys Asp	Phe Gly 340 Arg Trp Tyr Phe Leu 420 Thr Leu Ser Ser	Phe 325 Arg Asn Gly Leu His 405 Cys Ala Ser Lys Ala 485	Asn Leu Phe Phe 390 Cys Cys Thr Ser 470 Ser	Asn Gly Arg Ala 375 Ser Ala Gly Asn Ser 455 Ser	Ile Lys Ser 360 Phe Ile Met Arg Ile 440 Ser Ser Asp	Ala Arg 345 Val Phe Phe Lys Phe 405 Ile Thr Lys	Met 330 Ser Val Ala Asn Glu 410 Arg Lys Gly Thr Ser 490	315 Phe Asn Ser Trp Ser 395 Asn Leu Lys Ser Tyr 475 Leu	Cys Ile Arg Leu Gly 380 Leu Val Ala Ser Asn 460 Phe Ser	Val Thr Thr 365 Pro Gln Gln Asp Ser 445 Ser Lys Lys	Val Leu 350 Phe Leu Gly Lys Asn 430 Asp Thr Arg Leu	Met 335 Arg Leu Asn Leu Gln 415 Ser Asn Tyr Asn Ala 495	320 Val Glu Leu Ile Phe 400 Trp Asp Leu Leu Ser 480 His
Gly Glu Gly Pro 385 Ile Arg Trp Gly Thr 465 His	Ile Val Ile Val Met 370 Phe Arg Ser Lys 450 Ser Thr Asp	Met Cys Leu 355 Thr Met Ile His Lys 435 Ser Lys Asp Gly	Phe Gly 340 Arg Trp Tyr Phe Leu 420 Thr Leu Ser Ser Asp 500	Phe 325 Arg Asn Gly Leu His 405 Cys Ala Ser Lys Ala 485 Gln	Asn Leu Phe Phe 390 Cys Cys Thr Ser Ser 470 Ser	Asn Gly Arg Ala 375 Ser Ala Gly Asn Ser 455 Ser Met Ser	Ile Lys Ser 360 Phe Ile Met Arg Ile 440 Ser Ser Asp	Ala Arg 345 Val Phe Phe Lys Phe 405 Ile Thr Lys Ile 505	Met 330 Ser Val Ala Asn Glu 410 Arg Lys Gly Thr Ser 490 Pro	Asn Ser Trp Ser 395 Asn Leu Lys Ser Tyr 475 Leu Val	Cys Ile Arg Leu Gly 380 Leu Val Ala Ser Asn 460 Phe Ser His	Val Thr Thr 365 Pro Gln Gln Asp Ser 445 Ser Lys Lys Gln	Val Leu 350 Phe Leu Gly Lys Asn 430 Asp Thr Arg Leu Val 510	Met 335 Arg Leu Asn Leu Gln 415 Ser Asn Tyr Asn	320 Val Glu Leu Ile Phe 400 Trp Asp Leu Leu Ser 480 His

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aacaaaac														240
atggcgtg	***						-							300
atcaaaca														360 360
aaaaacaa	-			-										420
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gagaaatt														იიი ინმ
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tacattct		_												900
ctagcgag	_			-										960
gatgaatt				_	_									1020
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1		5					10			_		15		
Pro Pro		Leu	Glu	Asn	Leu		Pro	Glu	Asp	Ser		Leu	Val	
	20					25					30		_	
Arg Arg		Phe	Thr	Phe		Asn	Lys	Thr	Gly		Phe	Gin	Asp	
	35				40					45		_	_	
Val Gly	Pro Gln	Arg	Lys		Leu	Val	Ser	Tyr		Met	Ala	Cys	Ser	
50				55		_		_	60		~ 1	- 3		
Ile Gly .	Asn Ile	Thr		Gln	Asn	Leu	ГЛS	_	Pro	Val	GIn	тте		
65			70					75	_				80	
Ile Lys	His Thr		Thr	Gln	Glu	Val		His	Pro	He	Cys		Pne	
	_	85	_	_	_	,	90	~ -	_		en.)	95	G.1	
Trp Asp	Leu Asn 100		Asn	Lys	Ser	Phe 105	Gly	Gly	Trp	Asn	Thr 110	Ser	GIA	

	Cys	Val	Ala 115	His	Arg	Asp	Ser	Asp 120	Ala	Ser	Glu	Thr	Val 125	Сув	Leu	Суѕ
	Asn	His	Phe	Thr	His	Phe	Gly 135	Val	L∙eu	Met	Asp	Leu 140	Pro	Arg	Ser	Ala
	Ser 145	Gln	Leu	Asp	Ala	Arg 150	Asn	Thr	Lys	Val	Leu 155	Thr	Phe	Ile	Ser	Туг 160
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	Met	Asn	Leu 195		Thr	Ala	Leu	Leu 200	Phe	Leu	Asn	Leu	Leu 205	Phe	Leu	L∙∋u
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	Ala 235		Leu	L∙eu	His	Phe 230		Leu	Leu	Ala	Thr 235		Thr	Тгр	Met	Gly 240
		Glu	Ala	Ile	His 2:5	Met	Tyr	Ile	Ala	Leu 250		Lys	Val	Phe	Asn 255	Thr
	Tyra	Ile	Arg	Arg 250	Tyr	Ile	Leu	Lys	Phe 265	Сув	Ile	Ile	Gly	Trp 270	Gly	Leu
	Pro	Ala	Leu 275	Val	Val	Ser	Val	Val 230	Lesu	Ala	Ser	Arg	Asn 285	Asn	Asn	Glu
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L€				Asn	885					890					895	
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           IMAGE:84101 5', mRNA sequence.
ACCESSION
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VERSION
           T71087.1 GI:685608
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REFERENCE
           1 (bases 1 to 520)
 AUTHORS
           Hillier,L , Lennon,G., Becker,M., Bonaldo,M.F., Chiapelli,B.,
           Chissoe, S., Dietrich, N., DuBuque, T., Favello, A., Gish, W., Hawkins, M., Hultman, M., Kucaba, T., Lacy, M., Le, M., Le, N., Mardis, E., Moore
            ,B., Morris,M., Parsons,J., Prange,C., Rifkin,L., Rohlfing,T.,
            Schellenberg, K., Soares, M.B., Tan, F., Thierry-Meg, J., Trevaskis, E.,
           Underwood, K., Wohldmann, P., Waterston, R., Wilson, R. and Marra, M. Generation and analysis of 280,000 human expressed sequence tags
  TITLE
           Genome Res. 6 (9), 807-828 (1996)
  JOURNAL
  MEDITNE
           97044478
COMMENT
           Contact: Wilson RK
           Washington University School of Medicine
           4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
           Tel: 314 286 1800
           Fax: 314 286 1810
           Email: est@watson.wustl edu
           Insert Size: 592
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           clone is available royalty-free through LLNL; contact the IMAGE
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VERSION
KEYWORDS
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REFERENCE
           1 (bases 1 to 472)
 AUTHORS
           Marra, M., Hillier, L., Allen, M., Bowles, M., Dietrich, N., Dubuque, T.,
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  TITLE
           The WashU-HHMI Mouse EST Project
  JOURNAL
           Unpublished (1996)
COMMENT
           Contact: Marra M/Mouse EST Project
           WashU-HHMI Mouse EST Project
           Washington University School of MedicineP
            4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
           Tel: 314 286 1800
           Fax: 314 286 1810
           Email: mouseest@watson.wustl.edu
           This clone is available royalty free through LLNL; contact the
            IMAGE Consortium (infosimage.llnl.gov) for further information.
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                    3']; double-stranded cDNA was ligated to Eco RI adaptors
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ORIGIN

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                                                   linear
                                                           EST 01-MAR-1995
DEFINITION yc50d03.rl Stratagene liver (#937224) Homo sapiens cDNA clone
           IMAGE:84101 5', mRNA sequence.
ACCESSION
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VERSION
KEYWORDS
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SOURCE
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 ORGANISM Homo sapiens
           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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REFERENCE
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           Chissoe, S., Dietrich, N., DuBuque, T., Favello, A., Gish, W., Hawkins, M., Hultman, M., Kucaba, T., Lacy, M., Le, M., Le, N., Mardis, E., Moore
           ,B., Morris,M., Parsons,J., Prange,C., Rifkin,L., Rohlfing,T.,
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           Underwood, K., Wohldmann, P., Waterston, R., Wilson, R. and Marra, M.
           Generation and analysis of 280,000 human expressed sequence tags
 TITLE
 JOURNAL.
           Genome Res. 6 (9), 807-828 (1996)
 MEDLINE
           97044478
           Contact: Wilson RK
COMMENT
           Washington University School of Medicine
           4444 Forest Park Parkway, Box 8501, St Louis, MO 63108
           Tel: 314 286 1800
           Fax: 314 286 1810
           Email: est@watson.wustl.edu
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           clone is available royalty-free through LLNL; contact the IMAGE
           Consortium (info@image.llnl.gov) for further information.
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; Site_2: XhoI; Cloned unidirectionally. Primer: Oligo dT. Hepatectomy from normal male caucasian. Average insert size: 1.1 kb; Uni-ZAP XR Vector; ~5' adaptor sequence 5' GAATTCGGCACGAG 3' ~3' adaptor sequence: 5' CTCGAGTTTTTTTTTTTTTTTTTT 3'"

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Qу	2023	acaactcggaactt	ggctctcag	gogtato	catecetgt	taccagg	gacaaatgo	caatttca	2082	
•			11 H H H	ĪŦĪĦĿ	шшш					
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